

RAW SEQUENCE LISTING

DATE: 07/30/2001

PATENT APPLICATION: US/09/485,421

TIME: 16:07:28

Input Set : A:\Upap0350.app

Output Set: N:\CRF3\07302001\I485421.raw

3 <110> APPLICANT: Ayyavoo, Velpandi
 4 Patel, Mamata
 5 Kieber-Emmons, Thomas
 6 Weiner, David B.
 7 Mahalingam, Sundaramy
 9 <120> TITLE OF INVENTION: Functional Fragments of HIV-1 VPR Protein and Methods
 10 of Using the Same
 12 <130> FILE REFERENCE: UPAP0350
 14 <140> CURRENT APPLICATION NUMBER: 09/485,421
 C--> 15 <141> CURRENT FILING DATE: 2000-10-05
 17 <150> PRIOR APPLICATION NUMBER: 60/055,754
 18 <151> PRIOR FILING DATE: 1997-08-14
 20 <160> NUMBER OF SEQ ID NOS: 18
 22 <170> SOFTWARE: PatentIn Ver. 2.1
 24 <210> SEQ ID NO: 1
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 36 Asp Trp Thr Leu Glu Leu Leu Glu Glu Leu Lys Asn Glu Ala Val Arg
 37 20 25 30
 39 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
 40 35 40 45
 42 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
 43 50 55 60
 45 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
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 67 Lys His Phe Asp Pro Arg Leu Leu Thr Ala Leu Gly Asn His Ile Tyr
 68 35 40 45
 70 Asn Arg His Gly Asp Thr Leu Glu Gly Ala Gly Glu Leu Ile Arg Ile

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71          50          55          60
73 Leu Gln Arg Ala Leu Phe Met His Phe Arg Gly Gly Cys Ile His Ser
74 65          70          75          80
76 Arg Ile Gly Gln Pro Gly Gly Gly Asn Pro Leu Ser Ala Ile Pro Pro
77          85          90          95
79 Ser Arg Ser Met Leu
80          100
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93 1          5          10          15
95 Thr Ile Glu Glu Ala Phe Asp Trp Leu Asp Arg Thr Val Glu Ala Ile
96          20          25          30
98 Asn Arg Glu Ala Val Asn His Leu Pro Arg Glu Leu Ile Phe Gln Val
99          35          40          45
101 Trp Gln Arg Ser Trp Arg Tyr Trp His Asp Glu Gln Gly Met Ser Arg
102          50          55          60
104 Ser Tyr Thr Lys Tyr Arg Tyr Leu Cys Leu Met Gln Lys Ala Val Phe
105 65          70          75          80
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127          20          25          30
129 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
130          35          40          45
132 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
133          50          55          60
135 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
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155   20               25               30
157 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
158   35               40               45
160 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
161   50               55               60
163 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
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183   20               25               30
185 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
186   35               40               45
188 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
189   50               55               60
191 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
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211   20               25               30
213 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
214   35               40               45

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216 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
217      50                      55                      60
219 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
220 65                      70                      75                      80
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223      85                      90                      95
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231 <212> TYPE: PRT
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239 1      5                      10                      15
241 Asp Trp Thr Leu Glu Leu Leu Glu Glu Leu Lys Asn Glu Leu Val Arg
242      20                      25                      30
244 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
245      35                      40                      45
247 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu
248      50                      55                      60
250 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
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254      85                      90                      95
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270      20                      25                      30
272 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu
273      35                      40                      45
275 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Pro Leu Ile Arg Ile Leu
276      50                      55                      60
278 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg
279 65                      70                      75                      80
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291 <223> OTHER INFORMATION: Description of Artificial Sequence: Novel Sequence

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298 20 25 30

300 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu

301 35 40 45

303 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ser Leu

304 50 55 60

306 Gln Gln Leu Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg

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328 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu

329 35 40 45

331 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu

332 50 55 60

334 Gln Gln Ser Leu Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg

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356 His Phe Pro Arg Ile Trp Leu His Ser Leu Gly Gln His Ile Tyr Glu

357 35 40 45

359 Thr Tyr Gly Asp Thr Trp Thr Gly Val Glu Ala Leu Ile Arg Ile Leu

360 50 55 60

362 Gln Gln Leu Ser Phe Ile His Phe Arg Ile Gly Cys Arg His Ser Arg

VERIFICATION SUMMARY

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TIME: 16:07:29

Input Set : A:\Upap0350.app

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